



SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Stuiver, Maarten
 Ponstein, Anne
 Ohl, Stephan
 Goddijn, Oscar
 Simons, Lambertus
 Dekker, Bernardus
 Hoekstra, Sietske
 Tigelaar, Hendrik
 - (ii) TITLE OF INVENTION: Plasmids For Plant Transformation And Method For Using The Same
 - (iii) NUMBER OF SEQUENCES: 19
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Zeneca Ag Products Inc.

Elzinga, Nicolas

- (B) STREET: 1800 Concord Pike
- (C) CITY: Wilmington
- (D) STATE: DE
- (E) COUNTRY: USA
- (F) ZIP: 19850
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 09/469,812
 - (B) FILING DATE: 22-DEC-1999
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: EP 97201990.5
 - (B) FILING DATE: 30-JUN-1997
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT/EP98/04171
 - (B) FILING DATE: 29-JUN-1998
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Hohenschutz, Liza D.
 - (B) REGISTRATION NUMBER: 33,712
 - (C) REFERENCE/DOCKET NUMBER: MOG 57688/UST
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (302) 886-1699
- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA

(iii)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 1:	
GATGGCAA	AC GCTAATAAGG G	21
(2) INFO	RMATION FOR SEQ ID NO: 2:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
GAGCGGGT	TT AACCTACTTC C	21
(2) INFO	RMATION FOR SEQ ID NO: 3:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
CTGTTGTC	GA CGGCTGGCTG GTGGCAGGAT ATATTGTGG	39
(2) INFO	RMATION FOR SEQ ID NO: 4:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 38 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	

(iii) HYPOTHETICAL: NO

(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
CCAGCCGT	CG ACAACAGCTC CCCGACCGGC AGCTCGGC	38
(2) INFO	DRMATION FOR SEQ ID NO: 5:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
TCGACGGG	GT GCCTGGGGCG GTGGCGGGGC GGGCCGGGGC GTGC	44
(2) INFO	RMATION FOR SEQ ID NO: 6:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
TCGAGCAC	GC CCCGGCCCGC CCCCACGC CCCCAGGCAC CCCG	44
(2) INFO	RMATION FOR SEQ ID NO: 7:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	
(×i)	SEQUENCE DESCRIPTION: SEQ ID NO: 7:	
TTCTCGAG	AG GGTCGGTTAC TCG	23
(2) INFO	RMATION FOR SEQ ID NO: 8:	

	(A) LENGTH: 23 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ'ID NO: 8:	
TTGTCGA	CTC CAAGACGCCG AGC	23
(2) INF	ORMATION FOR SEQ ID NO: 9:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:	
AATTCAG	ATC TCCATGGATC GATGAGCT	28
(2) INF	ORMATION FOR SEQ ID NO: 10:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:	
CATCGAT	CCA TGGAGATCTG	20
(2) INF	ORMATION FOR SEQ ID NO: 11:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	

(i) SEQUENCE CHARACTERISTICS:

(iii) HYPOTHETICAL: NO

(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 11:	
GATCGGTC	GA CC	12
(2) INFO	RMATION FOR SEQ ID NO: 12:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 12:	
GACGCTAA	AG GCAAACTTGA TTC	23
(2) INFO	RMATION FOR SEQ ID NO: 13:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 13:	
TTAACCTA	CT TCCTTTGGTT CCG	23
(2) INFO	RMATION FOR SEQ ID NO: 14:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 14:	
AGCGGCCCA	AT TTTAGTAGAC GGG	23
(2) INFO	RMATION FOR SEQ ID NO: 15:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	

(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 15:	
ATCAATTC	GA ACCCCAGAGT CCCGTCCA	28
	RMATION FOR SEQ ID NO: 16:	
(1)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 16:	
CTGGATCGT	IT TCGCATGATT G	21
(2) INFOR	RMATION FOR SEQ ID NO: 17:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 17:	
CCAACTTAI	TC AGTGATAAAG AATCCGC	27
(2) INFOR	RMATION FOR SEQ ID NO: 18:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 18:	

27

(D) TOPOLOGY: linear

GAGATCAGAT TGTCGTTTCC CGCCTTC

(2) INFORMATION FOR SEQ ID NO:19:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

TNCAATTGAA AY

12